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SEQ ID 9850

MVKQVFPKGFLWGGATAANQCEGAYNVDGRGLANVDVVPTGEDRFALISGQKKMFDFEEGYFYPAKESIDFYHHYKEDLALLAEMGFKTYRMSIAW
TRIFPKGDELYPNEAGLQFYENIFKECRKYGIEPLVTITHFDCPIYLIKHYGGWRSRKMIGFYERLVRALFTRFKGLVKYWLTFNBINMILHAPFM
GAGLYFEDGENQEQIKYQAAHHELVASAIAVKIAHEVDPNNQIGCMLAAGQYYPNTCHPQDYWASMQKNRENYFFIDVQARGKYPNYAKKHFEHLG
ISIQMTAEDLALLRDYTVDFISFSYYSSRVASGNPTVSEQVQENIFASLKNPYLKSSEWGWQIDPLGLRITLNAIWDRYQKPMFIVENGLGAVDIP
DENGYVEDDYRIDYLRQHIAAMRDAIYVDGVNLIGYTTWGCIDLVSAGTGEMEKRYGFIYVDRNNKGEGTLKRYKKKSFYWYKKVIASNGSQIE

SEQ ID 9851

SEO ID 9852

MDGTFLRSDKTYDKARFSSLLTLMEKYDIKFVAASGNLYDQLLLNFLEYPNRIAYVAENGGRVIDQDGTLLKETYLSNDTVAAVLSYLYQNYPBTL ISLSGEKRSYLERRTPINRRTELEYYMPNFIYKDHLLPLDDDRYFQMTLWVNENLVSEMLLDISEHFKNHHIRLTSSGFGCIDVLPADVNKADGIA ILLEKWGLKQDQVMVFGDGGNDVEMLRAANISYAMSNAPEBIKAIAKYQTVSNDQDGVLETIENFLHKMSDNE

SEQ ID 9853

SEO ID 9854

mkktsslfsipymawlflfvlapvaliawnsffdinghftlanyqtffssgtylkmsfnsvlyagivsfitllisypaaylltklkhkqlwlmlvi lptwinlllkayafmgifgqqgginafltfigigpkqilftdfsflfvaayiblpfmllpibnalddidqnliyasddlganawqtfqkvifplbl ngvragvqsvfipslslfmltrliggnrvitlgtaibqhflitqnkgmgstigvililvmvaimwltkerka

SEQ ID 9855

SEO ID 9856

MLEQLGNPQNNLSAIHVVGTNGKGSTTSYLQHIFTNSGYQVGTFT8PYIVDFRERISIDGQMIPESDFIKLVETVRPVVERLHLETNLBPATEFEV ITVLMFYYFGNSCPVDIVIIBAGMGGYYDSTNMFKALAVTCPSIGLDHQEVLGRTYVDIAEQKVGVLKKGVPFVYANDRQDVEBVFQIKAKETHSQ TYRLHNDFYIKBEENYFNYIGPQANIDHIQLQMPGHHQVSNASIAITTSLLLRDKYPKLTLQTIKDGLEMTKWVGRTELIFPNVMIDGAHNNESVD ALVQVIKKYQQKNVHILFAAINTKPIESMLESLSSIAPVSVTSFDYPKSINLDKYPKAYTRV8D#KKWLHDINLTSDKDFYVITGSLYFISQVRQE LLLIKTTTT

SEO ID 9857

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RN - 440183-53-9 REGISTRY

FS - PROTEIN SEQUENCE

SQL - 478

PATENT ANNOTATIONS (PNTE):

Sequence | Patent

Source |Reference

Not Given|WO2002034771

|claimed SEQID

19850

SEQ	1	MVKQVFPKGF	LWGGATAANQ	CEGAYNVDGR	GLANVDVVPT	GEDRFAIISG
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51 QKKMFDFEEG YFYPAKESID FYHHYKEDLA LLAEMGFKTY RMSIAWTRIF

101 PKGDELYPNE AGLQFYENIF KECRKYGIEP LVTITHFDCP IYLIKHYGGW

151 RSRKMIGFYE RLVRALFTRF KGLVKYWLTF NEINMILHAP FMGAGLYFED

201 GENOEQIKYO AAHHELVASA IAVKIAHEVD PNNOIGCMLA AĞOYYPNICH

251 PODYWASMOK NRENYFFIDY OARGKYPNYA KKHFEHLGIS IQMTAEDLAL

301 LRDYTVDFIS FSYYSSRVAS GNPTVSEQVQ ENIFASLKNP YLKSSEWGWQ

351 IDPLGLRITL NAIWDRYQKP MFIVENGLGA VDIPDENGYV EDDYRIDYLR

401 QHIAAMRDAI YVDGVNLIGY TTWGCIDLVS AGTGEMEKRY GFIYVDRNNK

451 GEGTLKRYKK KSFYWYKKVI ASNGSQIE

HITS AT: 244-276

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